HABC Gentoype Data Submission Form

Submitting	Thibe dentoype but submission form
Investigator	
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Phone	415/885-7397
Lab Contact	Dr. Christian Vaisse
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Phone	
Date	5/18/2005
Submitted	
Polymorphism	
Information	
Name	3' UTR insertion variant
RS# or Unique	N/A
Identifier	
Gene	Leptin Receptor (<i>LEPR</i>)
Chromosome	chr1:65814779
position	
Alleles	-/cttta
Assay	
Information	
Genotyping	Template-directed dye-terminator incorporation assay with fluorescence polarization
Method	detection (FP-TDI). Reference: Hsu TM, et al. <i>Biotechniques</i> . 2001 Sep;31(3):560-8
Genotypes in	Yes. Caucasian genotypes were 1106: 513: 82 (del/del: del/ins: ins/del). African
Genotypes in	
HWE (Y/N)	American genotypes were 625: 499: 90 (del/del: del/ins: ins/del).
HWE (Y/N) (attach HWE	American genotypes were 625: 499: 90 (del/del: del/ins: ins/del).
HWE (Y/N) (attach HWE Form)	
HWE (Y/N) (attach HWE Form) Amount of	American genotypes were 625: 499: 90 (del/del: del/ins: ins/del). 5 ng/assay
HWE (Y/N) (attach HWE Form)	5 ng/assay
HWE (Y/N) (attach HWE Form) Amount of DNA used	5 ng/assay GGACAGTTGCTCACACTTTGTAG
HWE (Y/N) (attach HWE Form) Amount of	5 ng/assay
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTCTCTCCC
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTCTCTCTCC 10 x Platinum Taq buffer (Invitrogen) 0.5 μL
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR Components	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTTCT
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR Components and	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTTCT
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR Components and Concentrations	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTTCT
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR Components and	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTTCT
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR Components and Concentrations (Taq, buffer,	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTTCT
HWE (Y/N) (attach HWE Form) Amount of DNA used PCR primers PCR Components and Concentrations (Taq, buffer, MgCl ₂ ,	5 ng/assay GGACAGTTGCTCACACTTTGTAG GACTCTGGTTTCTTTTCT

	PCR program:
	avenue de la companya
	SNPPCR45
	95 for 2:00min
	92 for 10s
	58 for 20s
	68 for 30s
	GoTo 2 44 Times
	68 for 10:00min
	4 Forever
DOD G 1	end
PCR Cycling	
conditions	PCR cleanup program (SAP)
(time and	27.5 45 :
temperature	37 for 45min
for each step,	90 for 15min
# of cycles,	4 forever
etc.)	end
	TDI programs for 20 gyalog which is standard. For some assays we may use more or
	TDI program: for 20 cycles which is standard. For some assays we may use more or less cycles, between 5 and 80, for optimum result
	less cycles, between 3 and 80, for optimum result
	TDI_20
	95 for 2.00min
	95 for 15s
	55 for 30s
	Go to 2 19 times
	4 forever
	end
Detection	TTGTGTTATAATGGGTAATAAAGTGTAATAGATTA
Oligo(s)	
(if applicable)	
Other	The genotyping method was FP-TDI. After the PCR, the reaction was treated with
Reaction	shrimp alkaline phosphatase (SAP) for 90 minutes at 37°C. Subsequently, an
Conditions	appropriate dye terminator reaction was then initiated, containing 2 µL of 10 x
(detection	reaction buffer, 1 μL of the dye terminators (in this case, C/T), 0.05 μL of the
reaction	detection oligo (10 mM), 0.05 μL of Acyclopol, and 9.9 μL of ddH ₂ O. The samples
components,	were subjected to the extension reaction (TDI program as noted above) and
incubation	genotypes were read in a plate reader.
conditions, gel	
%, etc.)	
Other Assay	
Info	